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Timestamp: [year=2008; month=10; day=20; hr=12; min=48; sec=14; ms=664;
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Application No: 10526049 Version No: 2.0

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Output Set:

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<140> 10526049

<141> 2008-09-17

<150> PCT/JP2003/010541

<151> 2003-08-20

<150> JP 2002/253752

<151> 2002-08-30

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gttcgagaga aggaaac atg cac aac gac aac act ccc cac tcg cgt cgc 290

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15 20 25

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30 35 40

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Arg Ala Leu Ala Asp Asn Pro Gly Thr Ala Pro Leu Asp Thr Phe Met

45 50 55

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		80		85		90	
gac agc ctg ccg cag ctc gcc ggc gcg ctc gcg tcc ggt tcg ctg acg	578						
Asp Ser Leu Pro Gln Leu Ala Gly Ala Leu Ala Ser Gly Ser Leu Thr							
		95		100		105	
cct gaa cag gaa tcg ctc gca ctg acg atc ctc gag gcc tgg tat ctc	626						
Pro Glu Gln Glu Ser Leu Ala Leu Thr Ile Leu Glu Ala Trp Tyr Leu							
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Gly Ile Val Asp Asn Val Val Ile Thr Tyr Glu Glu Ala Leu Met Phe							
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ccc ggc ttc tgg gcc gac aaa ccg atc gag agg caa gcc tg atg gcc	769						
Pro Gly Phe Trp Ala Asp Lys Pro Ile Glu Arg Gln Ala Met Ala							
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gat acc gat acg caa aag gcc gac gtc gtc gtc gtt gga tcg ggt gtc	817						
Asp Thr Asp Thr Gln Lys Ala Asp Val Val Val Val Gly Ser Gly Val							
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Ile Leu Leu Glu Ala Gly Pro Arg Met Pro Arg Trp Glu Ile Val Glu							
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cgc ttc cgc aat cag ccc gac aag atg gac ttc atg gcg ccg tac ccg	961						
Arg Phe Arg Asn Gln Pro Asp Lys Met Asp Phe Met Ala Pro Tyr Pro							
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Ser Ser Pro Trp Ala Pro His Pro Glu Tyr Gly Pro Pro Asn Asp Tyr							
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ctg atc ctg aag ggc gag cac aag ttc aac tcg cag tac atc cgc gcg	1057						
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Pro Ile Gln Tyr Asp Asp Leu Glu Pro Tyr Tyr Gln Arg Ala Glu Glu							
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Arg Lys Gln Pro Tyr Pro Met Pro Pro Leu Pro Leu Ser Phe Asn Glu							
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Gln Thr Ile Lys Thr Ala Leu Asn Asn Tyr Asp Pro Lys Phe His Val							
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Val Thr Glu Pro Val Ala Arg Asn Ser Arg Pro Tyr Asp Gly Arg Pro							

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Thr Cys Cys Gly Asn Asn Asn Cys Met Pro Ile Cys Pro Ile Gly Ala			
380	385	390	
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Met Tyr Asn Gly Ile Val His Val Glu Lys Ala Glu Arg Ala Gly Ala			
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aag ctg atc gag aac gcg gtc gtc tac aag ctc gag acg ggc ccg gac			1537
Lys Leu Ile Glu Asn Ala Val Val Tyr Lys Leu Glu Thr Gly Pro Asp			
415	420	425	
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Pro Lys Ile Leu Leu Met Ser Ala Asn Arg Asp Phe Pro Asn Gly Val			
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Ala Asn Ser Ser Asp Met Val Gly Arg Asn Leu Met Asp His Pro Gly			
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acc ggc gtg tcg ttc tat gcg agc gag aag ctg tgg ccg ggc cgc ggc			1777
Thr Gly Val Ser Phe Tyr Ala Ser Glu Lys Leu Trp Pro Gly Arg Gly			
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Pro Gln Glu Met Thr Ser Leu Ile Gly Phe Arg Asp Gly Pro Phe Arg			
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Ala Thr Glu Ala Ala Lys Lys Ile His Leu Ser Asn Leu Ser Arg Ile			
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gac cag gag acg cag aag atc ttc aag gcc ggc aag ctg atg aag ccc			1921
Asp Gln Glu Thr Gln Lys Ile Phe Lys Ala Gly Lys Leu Met Lys Pro			
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Asp Glu Leu Asp Ala Gln Ile Arg Asp Arg Ser Ala Arg Tyr Val Gln			
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Phe Asp Cys Phe His Glu Ile Leu Pro Gln Pro Glu Asn Arg Ile Val			
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Pro Ser Lys Thr Ala Thr Asp Ala Ile Gly Ile Pro Arg Pro Glu Ile			
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Thr Tyr Ala Ile Asp Asp Tyr Val Lys Arg Gly Ala Ala His Thr Arg			
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Glu Val Tyr Ala Thr Ala Ala Lys Val Leu Gly Gly Thr Asp Val Val			
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Phe Asn Asp Glu Phe Ala Pro Asn Asn His Ile Thr Gly Ser Thr Ile			
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acc gcc atg ccg gta ccg atg ctc gcc aag atc tac acg agc aac atc	2547		
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765	770	775	
ggc gcc aag ccg tac gcg gcc gcc ctt gcc gcc atc gcc aaa tgg acg	2643		
Gly Gly Lys Pro Tyr Ala Gly Gly Leu Gly Gly Ile Gly Lys Trp Thr			
780	785	790	
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Pro Val Lys Gln Ala Pro Pro Lys Asn Glu Ile Pro Ala Leu Leu Ser			
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Met Arg Trp Pro Leu Lys Ile Trp Asn Trp Leu Phe Leu Lys Asp Gly			
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ccg tac cag ccg aag ccg tcg cag agc gcc gaa tgg aat cgc gcc gcg	2931		
Pro Tyr Gln Pro Lys Pro Ser Gln Ser Ala Glu Trp Asn Arg Gly Ala			
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Tyr Leu Val Gln Gly Leu Ala His Cys Ser Thr Cys His Thr Pro Arg			
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ctc gcg ggg tcg gtg ctc gcc gcc tgg gac gcc tac aac atc acg tcg	3075		
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Asp Pro Asn Ala Gly Ile Gly Ser Trp Thr Gln Gln Gln Leu Val Gln			
940	945	950	
tat ttg cgc acc gcc agc gtg ccg gcc gtc gcg cag gcg gcc ggg ccg	3171		
Tyr Leu Arg Thr Gly Ser Val Pro Gly Val Ala Gln Ala Ala Gly Pro			
955	960	965	
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Ser Asn Ala Lys Gln Pro Arg Ser Ser Trp Gly Lys Pro Ala Glu Asp				
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Gln Ala Leu Gln Lys Gly Ser Phe Lys Thr Ala Asp Ser Leu Pro Gln			
	85	90	95
Leu Ala Gly Ala Leu Ala Ser Gly Ser Leu Thr Pro Glu Gln Glu Ser			
	100	105	110
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	115	120	125
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 50 55 60
 Tyr Pro Ser Ser Pro Trp Ala Pro His Pro Glu Tyr Gly Pro Pro Asn
 65 70 75 80
 Asp Tyr Leu Ile Leu Lys Gly Glu His Lys Phe Asn Ser Gln Tyr Ile
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 Arg Ala Val Gly Gly Thr Thr Trp His Trp Ala Ala Ser Ala Trp Arg
 100 105 110
 Phe Ile Pro Asn Asp Phe Lys Met Lys Ser Val Tyr Gly Val Gly Arg
 115 120 125
 Asp Trp Pro Ile Gln Tyr Asp Asp Leu Glu Pro Tyr Tyr Gln Arg Ala
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